DATE: 07/24/2001 TIME: 15:36:40

OIPE

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M1-2,4-5
                               Input Set · D:\Pa_00349.rpt
                               Output. Set.: N:\CRF3\07242001\I849529.raw
  1 <110 > APPLICANT: Fincher, Karen L.
                 Ziegler, Todd E.
  4 :120 - TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
                Plants
  7 <130 - FILE REFERENCE: 38-21(51893)B
 9 <140> CURRENT APPLICATION NUMBER: US/09/849,529
 9 <141> CURRENT FILING DATE: 2001-05-07
 3 (150) PRIOR APPLICATION NUMBER: US 60/196,868
11 <151: PRIOR FILING DATE: 2000-05-09
13 (160) NUMBER OF SEQ ID NOS: 24076
15 <210 - SEQ ID NO: 1
16 (211) LENGTH: 592
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20 -02230 OTHER INFORMATION: Clone ID: LIB3543-001-Q1-K1-A1
17 -CLIL: TYPE: DNA
22 -: 400> SEQUENCE: 1
24 chaaqcaaqt tiggaqaagt totoaagogo tactaatooa gaigtigigg aacagagtot
26 quataqtitiq qotqaqaaqq caaacaaaqa qatqqootoq qoaqtqaata qotoqqooqa
                                                                                                                                      -120
28 qaaqatgaaq aacgaqatgg ccatgacagt aaatagctcc actgaggaga tgaaaagaga
                                                                                                                                        240
30 queaucttea ataaaqaget cageggagaa gatgaacaaa gagacageet tgaaaataaa
                                                                                                                                        300
31 taact.caget gagaagatga agaaagaaac atcattaaat ageteagetg agaagatgag
34 acaaqaqaca gotqoaqaaa atqqttcaqo tqaaaaqatq aaqaaaqaqa caqoottgac
                                                                                                                                        360
                                                                                                                                                  2.217. 22227.
36 agtaaatato toagotgaga agatgaacaa agagacagoo tigacaataa ataactoago
                                                                                                                                        420
38 tyaqaaqaty aacaaaqaqa tyyccayyac aytyaatayc tcayctyaga ayatyayaca
                                                                                                                                       480
                                                                                                                                        540
40 agagacaget acagtuaatg geteggetga gaagalgaag aaagagaegg cetligaeagt
                                                                                                                                        592
42 agagageted aacqadaagg tgaagaaaga aacageetty acagtaaata ac
45 <2:10> SEQ ID NO: 2
                                                                                                                                                       i gravela and
46 - . Fire LENGTH: 531
                                                                                        same che
47 <1112 TYPE: DNA
48 <213> ORGANISM: Gossypium hirsutum
50 <223 OTHER INFORMATION: Clone ID: LIB3543-001-Q1-K1-A10
52 -400 SEQUENCE: 2
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Se fracticetti fitalfacatq chaqtifiqda tocaqtitatc aqqicticatta afqtiafiliq
                                                                                                                                        120
58 gattitttit tittaaattig alaalligta callactiag atteelgaea litalattal
                                                                                                                                        1300
60 aatactitga ataigittaa aaaaaaaita aacgcattig aaaitattaa teigaigait
                                                                                                                                        240
62 occarcatto acceptateget acatategete thephograpot egocacttee catteephtea
                                                                                                                                        300
64 cattqtttta atattaatat tiittaattaa catcaaccit tiittgiaaaa ciitaaaiittg
66 thoaaaaato ttaaattitta aatiqtatti aatattitoo aftiilogtia caftiitagi
                                                                                                                                       420
                                                                                                                                       480
68 qtt.ttatttt ocacactgaa atcittitgot titigotgata aaatittocca aaafgqttag
                                                                                                                                       531
70 tattititoa aaaittodig aliittataa alalactaac ccaalagata ti
13 - 210 - SEQ ID NOT 3
        The state of the s
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RAW SEQUENCE LISTING

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```
82 conacquete eqeteaqait tictoloagi qittaatagi gaaacgatgi ccactglogg
84 ajaagotgot tgotottaog otgotottat totthatgan ganggoando coardadogo
86 tyaqaaqatt totacaotag ttaaaqotqo aaatqtotot qttqaatoot actggocaag 180
                                                                        240
83 citqtteqet aagettittg agaaqtgeaa cattgaggat cicataacca atgitggage
90 taqtaqtaqt qqtqccccaq taqctqcaqc tqcacctqtt qctqctqqcq qaqqaqccqq
                                                                        3:):)
92 tyckyctyct ccacctccty ctyaydaaaa gaaggaagaa ccyyaggaag agagtgatga
94 cgatatqqqa ttcaqtttqt ttqattaqqt qtqttttqqt caattactaq ttatatcaca
                                                                        4.20
96 tqaaqqtaqt aacqetttta qaqttqtatt qetattqqta tttqcatqtt qattaqaaat
                                                                        480
                                                                        540
98 attacqtttt tqttqaatga aagttqacac taagtagctc tttctttqtq atctgttttt
100 tttgaatett eetatatata gaagtattat agatteta
                                                                         578
103 <210 > SEQ ID NO: 4
104 <211 - LENGTH: 645
105 <212 - TYPE: DNA
106 <213 - ORGANISM: Gossypium hirsutum
108 -320 - FEATURE:
109 CLDIE NAME/KEY: unsure
110 + 222 + \text{LOCATION}; (1) ... (645)
111 -0.23: OTHER INFORMATION: unsure at all n locations
112 (223) OTHER INFORMATION: Clone ID: LIB3543-001-Q1-K1-A2
114 (400) SEQUENCE: 4
                                                                          ti()
116 eggaegegtg ggaataactt caatgggtge gtgaatttaa ggaagatact taatetttea
118 agagecatea acettgaata cetttattge agtgaetgta aaagtttagt tgaactttgg
                                                                         1.20
120 aatqqaqatq attqtqtqqa ttttqttaat ttaaqqqaaa ttaacttcaa tgggtqcgtg
                                                                         180
                                                                         240
122 aatttaagaa agatacttaa tettteaaga geeateaace tegaatacet ttattgeagt
124 quotiqtaaaa gittagitga actitiggaat ggagatgatt gigiggatti tittaattia
126 agggaaatta aetteaatgg gtgegtgaga ttaaggaagg aetttgttaa tttaaeatee
                                                                         360
128 figuageast traagaaaaa tagetaatet treaagagee areaacettg aatteatrga
130 ingenetaat ighaaaagte tygttgaact teeftgeete gateattigg catetetiga.
                                                                         480
132 tigographia tottoatottq acqqatqcta taqtotcaaq aaqttootca qqtoccacqt
                                                                         540
134 cacttttgtt atttagattt atcaaaaacc agaatagnaa atatgtcgtt gcctttcgat
                                                                         600
136 cocchatgta atctteagea ettagatatg agtggeteag eggte
                                                                         645
139 -2105 SEQ ID NO: 5
140 THIT (ENGTH: 607
                                           Adres iver de prevener parque
141 - 12125 TYPE: DNA
142 <213> ORGANISM: Gossypium hirsutum
144 <223 OTHER INFORMATION Clone ID: LIB3543-001-Q1-K1-A3
146 <400% SEQUENCE: 5
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                                                                          +;(1
150 aataggetty aaaaagaaty geaateaace attacaaagy geagcageta atgetaaaty
                                                                         100
                                                                         180
152 catcagatga totacatgga caaaatcagg gcaaccaacc aaagggaatg gcaaaaatca
                                                                         240
154 cageettete ecaactatae tytytätäea taaatteaet aataateaaa atettyttiä.
156 tytactyaca aattacacay ycaacttyce eccatttacy tittyattya agattyaane
                                                                         3(1()
                                                                         360
158 aaccaaccaa cotoqattta aaqacqqaaa qtaqtaqcaa aaatotcaqo qqatoqtaat
160 tgtcatttgc agatgcacac tttgattacg taaaaggaag tagtccttca aattgacact
                                                                         4110
                                                                        480
162 accegaaaqaa ttacatttat ccaacaaatc aacaaqcaaa qqtcacaqat qccatttcaq
                                                                         540
164 tgctaactig itgcctcaca itglicating cittingfic andiagadig coactetiti.
166 qtqcaatqtc accaaaqqqa qqtaaaatct aaaataqtcc ctaattqtat attitaacaq -600
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- 172 2115 LENGTH: 653 173 < 212 - TYPE: DNA 174 < 213 - ORGANISM: Gossyprum hirsutum $176 \times 220 \times \text{FEATURE}$ 1 7 (22) NAME/KEY: unsure $1^{\circ}3 = 222 + \text{LOCATION}: (1)..(653)$ 17) < 223 - OTHER INFORMATION: unsure at all n locations 180 <223 - OTHER INFORMATION Clone ID: LIB3543-001-Q1-K1-A4 182 <400 - SEQUENCE: 6 184 qtqaqqaaaa aaaagettca atetttggtt gttaagtqgg tgaaacaaaa aaaaaaatqg 186 aqqqcaaaqa qqaqqatqtt aaactaggag ccaacaaqtt ctctgagaga caacctattg 120 188 ggadateago temameagad maggadtada aggagddadd accagdadda ftgtftgagd 180 190 chgghgaget caugheatgg feithetaea gageeggeat tgetgagtti gifgehaeet 240 300 192 totligttoot thicateact glottgactg loatgggfgf gfofcaafof aagaccaaat 360 194 grantactiff tygitational agentificit gageotitigg tygicalgale liligeneily 420 106 fituitiqtae tqetqqeatt teaqqtqqee atattaaecc agetqtqaec fftqqqtfqt 480 198 tactugeday gaugetytee ettacaaggy cagtatteta catgateaty caatgeetty 540 200 qtqctatctq tqqaqctqqt qtaqtqaaag gattcgaagg agacagtaga tatqaqatgt 600 > 202 tggngtggtg gagccaatgt tgtgaaccat ggctacacca aagggtgatg gtcttggtgc 653 > 204 tqaaatcatt qcacttttqn tcttqnctac actgqtttct ctqctactga tgc 207 <210: SEQ ID NO: 7 208 <211 - LENGTH: 638 209 <212 TYPE: DNA 210 <213> ORGANISM: Gossyprum hirsutum 212 <220: FEATURE: 213 - 221 NAME/KEY: unsure 214 <222 LOCATION: (1)..(638) 215 (223) OTHER INFORMATION unsure at all n locations 216 - 223 - OTHER INFORMATION, Clone ID: LIB3543-001-Q1-K1-A5 218 <400 > SEQUENCE: 7 220 ottaattiggo tigogotottig ottigttogto toaatootigo tattitotata agaatagitti 60 222 aggitticat tilligagic aaactaicag titacticci ciaicatcgc aaaticigaa 120 180 ...: qqaanaqqqa tooqatetga tetyetqeaa atgggqqaga agtqqqqtqtt qatqaaaanq 240 226 gataagaaqa ataacaacat tigaagtigatt aagtattiggig iggaagaggiga tigagaacaatti 300 228 atttrgoogg temmegatag catemagegit metetigate etgetemiet etgemeeme 360 2.0 arcaecqteq etqttaqeec tqetttecaq caaqateqta tqtqqetcaa tqqeaaqqaa $z \cdot z$ aratheettt deggaggaag gladdagaad lyftlyaggg aauteagaaa uegggelagg z-4 qaaqifqaqq afaaaqaqaq qqqaataaaq afaqaqaaqa aaqattqqqa qaaacfqcaf 480 gase entroatatiq caretiataa caattteeet, acagetgeag gaftggette efetgetget 540 -> 238 ggttntgctt gtcttgtttt tgcacttgca aagcttatga atgtcagtga agatgagagt 600 4-> 240 caactttotg ctattgotag gcaaggttoa ngcagtgo 638 $244 < 210 > \mbox{SEQ}$ ID NO: 8
 - 244 ×211> LENGIH: 636
 - 245 <212> TYPE: DNA
 - 246 213 ORGANISM: Gossypium hirsutum
 - 24% 220% FFATURE:
 - , 4 * /, 1 NAME/KEY: dostro

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```
252 < 223 > OTHER INFORMATION: Clone ID: LIB3543-001-Q1-K1-A6
    254 - 400 > SEQUENCE: 8
    256 flottofloa aarigittit oodagitati tigittiggit tiolitigato tolototolo
    258 retetiticat ericaatett tittetyttet tittetititt aaaittatat aaaattaty
                                                                              120
    200) aatatattae giilliggile lyygityatt täyteäytti aeticetitt itaittataa.
                                                                              18:)
    2.62 tettititie at categorie teetitigiat tietaraaca titattacea titattiaat
                                                                              240
                                                                              3:)1)
    264 attitigaget talacacity attitegatt accellitigt gagallegaa aaaaaaaace
    266 cattittita ettättitee qaiqaqqaqa qqqaqqqqte tegeeqeeqe eqicateeee
                                                                               360
    268 gatggagaag aacccaaata cagaggtgtt eggaaaagan catggggtag attegeaget.
                                                                              420
    270 qaqatooqaq accoqtqqaa aaaaaccaqq qtttqqttaq qtactttoqa ttooqooqaa.
    272 gaageagete gagettaega taeggeggeg atgaeaetge gtggaeegaa ageeaaaaeg
                                                                              540
\psi_{t \, 	au} > \, 274 aattteecaa teaateeete gaataetane egettteega tegateacea eeacegtegg .
                                                                              600
    276 ttatacetga tgggtaattt teaagaeeeg aaaatg
                                                                              ń∃h
    279 <210 - SEQ ID NO: 9
    280 <211> LENGIH: 663
    281 - 212 - TYPE: DNA
    282 <213 : ORGANISM: Gossypium hirsutum
    284 <220. FEATURE:
    285 (221 - NAME/KEY: unsure
    286 <222 LOCATION: (1)..(663)
    287 < 223: OTHER INFORMATION: unsure at all n locations
    288 <223 > OTHER INFORMATION: Clone ID: LIB3543-001-Q1-K1-A7
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#_-> 292 ccccgcgtcc ggtgagnaaa tgaataatcc actaccgaca ctgtctttcg tgtaaagctt
    194 cagadatace attacagggt acagagattg agagaaacta cttgtgggad ttcacceccc
                                                                              120
                                                                              1.80
    19th coccasaaaaa chqttttttc totgtototg caataaactg gaaggagcag acacagagac
    298 agqqaqaqaqaqaqaqaqaq agaqaqaqaqaqaqaqaqaqaqqaqaqaqaqq
                                                                              240
    300 caqtatqqct aaqttaatca aaggataaga titgatccar cfcfcffafg ffffccffff.
                                                                              3000
     301 tettiqtata aaqteeaqaa gaageaqatt caaqqatatt qagaaqaagt ttetaaagat
    304 titightaage tagetatggg aactetigea alegalaegg agigigigga tietgalata.
                                                                              420
    306 tancqtqcit tqqqattcaa agaattgggt aaaggagttg ttggaactcc acggatttta
                                                                              480
     308 teachtectit efficients tigaggaatee gfeeaaagaa aegagafgga gfeegagaeg
                                                                              5.10
  > 310 gtgaacttca aanataatgt tgccatgttt catgggttaa gagctccacc ataagcattc
                                                                              600
     312 agoaftatat egategeate ticaagtaig eiggeigeag teegiegigt tittelegiig
                                                                              hh(I
                                                                              hh 3
    314 cat.
    fl^{2} < 210 > SEQ ID NO: 10
                                                         J.d. m.
     518 - 211 - LENGTH: 200
    319 <212 - TYPE: DNA
    3.0 - 213 - ORGANISM: Gossyp.um hirsutum:
    Ref. - 223 - OTHER INFORMATION: Clone ID: LIB3643-00第-y1-K1-A8
    32: -400 - SEQUENCE: 10
    326 tocatatoot, tygatygyttt atotoaatyg toatyctaat actititiytt ytacfoffic
    328 fetaggatte tifettitet tgaactecat tgicaaaagg etigtaccaa gitgfaafge
                                                                              110
    330 caqtattaat qtaqtcaata aaaactatat qtafqqtaaa qtatqtattt ttattafftq
                                                                              180
    332 citacitcae tygyttygee
                                                                              200
    3.65 (210) SEQ ID NO: 11
    KON THE LINGSHIP SEE
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                                                                          1.20
316 theagageea taatttaett ettgeggaga glyagggate atalaeggig eagcagaaet
                                                                          180
343 aractayett agacatacat qtaggycayt catafteytt tefyetaace aeggateaga
35)) atgruagtag egattaetae attgtageaa gtgeraggtt tgtgaatgaa tegeagtgga
                                                                          240
352 alogiqtiae egytyttyce attitgegit affetaatte yaaaygaaay geateegyte
                                                                          300
354 rectedectga tecacegoag gatgaattog acaaaacett ticaatgaac caagetagat
                                                                          4.20
356 ctatcagaty qaatqtatet gecagtggtg etegecegaa cecacaggga tegtttagat.
358 atyy!tegat taatgtgaet gaagtttaeg tettgagaaa caaaccacca gagactattg 480
                                                                          540
360 atgggaaaag acgagegaea eteagtggga tateetitigt eaateetiget acaccaatee -
362 gyettyttyä eeaytataay attäaayyay tatataaget tyä
365 <210 - SEQ ID NO: 12
366 <211 · LENGTH: 672
367 (212 - TYPE: DNA
363 (213) ORGANISM: Gossypium hirsutum
370 < 320 + FEATURE:
371 < 121 · NAME/KEY: unsure
372 < 222 < LOCATION: (1)...(672)
373 <223 OTHER INFORMATION: unsure at all n locations
374 <223 OTHER INFORMATION: Clone ID: LIB3543-001-Q1-K1-B10
376 <400 · SEQUENCE: 12
378 teaaqottea qeagaaggaa atatteggta aaaaqgagag attetgaagg agagagtagt
380 tectecacet ggggaaagaa gttgagtggg teateteegg agggaetgaa caatagttet
                                                                          120
381 ageotoagtig attictatito tigaticaagg ogagocagaa attigitotto taatagiggat
                                                                          180
384 tytyytätty ettetteagt taggaeteag agateaaaca gtagttatyy tagtyggaga
                                                                          240
                                                                          300
38n cheectaace aaqeaaatgg aaataqtita actitgaatg agteaceeat ggteateeca
                                                                          360
388 caggearche aarchgahat tegtachgah afgaafgete etgtheecah agaaactgee.
390 feaactegta ctagttetta cagtegatea ggtagtafag atgagagttt acgaggttte 420
392 algeetteea gleetteaga agttagtggt faccaclett cagligaacca ggglagette -
394 cagcactaca atatggatgg ctttgcagag gtattattag aacttgagag aattgaacaa
396 datquaqadt tgacatacga gcaattactt ditctagada ccagcitigci ccittaatggci
                                                                          600
                                                                          660
348 etggaetttt atgaategea tagagatatg agattggata tagatgatat gteatatnga
                                                                          672
400 quattactag ct
                               the state of property of many color than a second
40 + <210 > SEQ ID NO: 13
404 <211> LENGTH: 260
40% ARIZE TYPE: DNA
400 - 213 - ORGANISM: Gossypium hirsutum
408 - 223 - OTHER INFORMATION: Clone III: 1183543-001-Q1-K1-B11
410 - 400 - SEQUENCE: 13
41. atgregactor officetting officettic occupioned actiffingua forcagatoa
414 gyaaattato aaatoogago ogooatggag gitaacgaca agoiiitigaa caogggiatt
                                                                          1.20
416 chratagtag ccareetagt agtggeeaag chratatett techrataan geetagatet.
                                                                          180
418 auquaanggg ttocaceage aqteaaaace tqqccqqtaa tcqqtqqqct ccttcqatca
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420 tgaaaggeet ategtgatge
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423 - 210 - SEQ ID NO: 14
424 - 211 - LENGTH: 653
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L:9 M·271 C: Current Filing Date differs, Replaced Current Filing Date
L.134 M·341 W: (46) "n" or "Xaa" used, for SEQ ID# 4
L 202 M 341 W. (46) "n" or "Xaa" used, for SEQ ID# 6
L 204 M 341 W. (46) "n" or "Xaa" used, for SEQ ID# 6
L 238 M 341 W (46) "n" or "Xaa" used, for SEQ ID# 7
L-240 M-341 W (46) "n" or "Xaa" used, for SEQ ID#-7
L 274 M.341 W. (46) "n" or "Xaa" used, for SEQ ID#.8
L 292 M:341 W (46) "n" or "Xaa" used, for SEQ ID# ")
L-310 M-341 W. (46) "n" or "Xaa" used, for SEQ ID#:9
L 398 M.341 W (46) "n" or "Xaa" used, for SEQ ID#:12
I. 456 M 341 W (46) "n" or "Xaa" used, for SEQ ID#:14
L 490 M 341 W (4b) "n" or "Xaa" used, for SEQ ID#:15
L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
1..556 M.341 W. (46) "n" or "Xaa" used, for SEO ID#:17
I-622 M.341 W. (46) "n" or "Xaa" used, for SEO ID#:19
L:782 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#:24
L:838 M:341 W (46) "n" or "Xaa" used, for SEQ ID#:26
L.946\ M.341\ W:\ (46)\ "n"\ or\ "Xaa"\ used, for SEQ\ ID#:29
L:948 M.341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1084 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1116 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:1118 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:1152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:1154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:1172 M:341 W: (46) "n" or "Yaa" used, for SEQ ID#:36
L 1208 M:341 W. (46) "n" or "Xaa" used, for SEQ ID#:37
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L 1286 M.341 W (46) "n" or "Xaa" used, for SEO ID# 39
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L-1366 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L 1402 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
! 1404 M:341 W (46) "n" or "Xaa" used, for SEQ ID#:42
1..1420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#.43
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L:1524 M 341 W: (46) "n" or "Xaa" used, for SEQ ID# 46
1 17 m M.341 W (4m) "n" or "Kaa" used, for SEQ ID# 53
I 1733 M:341 W (4h) "h" or "Kai" used, for SEQ ID# 33
] [238 M:341 W (46) "n" or "%aa" used, for SEQ ID# 54
                 Tither "h" or "Raa" ased, for SEQ ID# of
1 1 1 m M: 341 W
                 (46) "n" or "%11" usel for SEQ ID# 56
  1833 M:341 W
L.1894 M:341 W (46) "n" or "Xaa" used, for SEQ ID#:57
I. 1896 M:341 W (46) "n" or "Xaa" used, for SEQ ID#.57
L:1964 M:341 W. (46) "n" or "Xaa" used, for SEQ ID# 59
I:1982 M:341 W: (4b) "n" or "Xaa" used, for SEQ ID#.60
1:2000 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:60
1 - 30 % M - 34% W: (46) "n" or "Xaa" used, for SEQ ID#:62
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L 2140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64
L:2162 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65

Volume in drive D:\ is 010504 0932 Directory of D:\

pa_00349.rpt

.9401 KB - 5 4 ...

1 file(s) Total filesize 19461 KB O kilobytes free

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